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1	Met Thr Leu Ile Tyr Val
61 7	cttcaatatttacaatggtcccctcaatcacacggattgtactggttaacattctgttgg Pro Scr Ilc Phe Thr Met Val Pro Ser Ile Thr Arg Ile Val Leu Val Asn Ile Leu Leu
121 27	cgacgttggttttgggagctgcagtccttccacgagacaacagaactgtttgcgggagtc Ala Thr Leu Val Leu Gly Ala Ala Val Leu Pro Arg Asp Asn Arg Thr Val Cys Gly Ser
181 47	aactctgcacatggtggcacgactccggcgagataaacaccggtactcctgtacaggcag Gin Leu Cys Thr Trp Trp His Asp Ser Gly Glu lie Asn Thr Gly Thr Pro Val Gin Ala
241 67	gaaacgttcgacaatcccgaaagtactctgtccatgtgagcctggcagaccgtaaccaat Gly Asn Val Arg Gln Ser Arg Lys Tyr Ser Val His Val Ser Leu Ala Asp Arg Asn Gln
	tctacgactctttcgtatatgaatcgatacctaggaacggcaatggcagaatttattctc Phe Tyr Asp Scr Phe Val Tyr Glu Scr Ilc Pro Arg Asp Gly Asp Gly Arg Ile Tyr Scr
361 107	ccaccgaccacctaacagcaatacattgaatagtagcattgacgacggtatatcaatcg Pro Thr Asp Pro Pro Asn Ser Asn Thr Leu Asn Ser Ser Ile Asp Asp Gly Ile Ser Ile
421 127	aaccatctctcggcatcaacatggcttggtcccagttcgaatatagacgagatgtcgaca Glu Pro Ser Leu Gly Ile Asn Met Ala Trp Ser Gln Phe Glu Tyr Arg Arg Asp Val Asp
	ttaagattactacaatcgatggctcaatattggatggccctttggacattgttattcggc IIc Lys IIe Thr Thr IIe Asp Gly Ser IIe Leu Asp Gly Pro Leu Asp IIe Val IIe Arg
	cgacttctgttaagtactcagtcaaaagatgtgtgggtgg
	atgateceaatggtegaaaattetetgttgagttaaagagtgaeetttaeagttaeetet Tyr Asp Pro Asn Gly Arg Lys Phe Ser Val Glu Leu Lys Ser Asp Leu Tyr Ser Tyr I
	ccgacggttcgcaatatgtgacctctggagggagcgtggttggt
	ccctggtgatctttgccagccctttcttgccacgggatatggttcctcatatgacaccac Ala Leu Val Ile Phe Ala Ser Pro Phe Leu Pro Arg Asp Met Val Pro His Met Thr Pro
	acgacacccagacaatgaagccgggcccaatcaataatggggactggggttcaaagccta His Asp Thr Gin Thr Met Lys Pro Giy Pro Ile Asn Asn Giy Asp Trp Giy Ser Lys Pro
	tactctacttcccgcctggcgtatactggatgaacgaggatacctctggtaaccccggga Ilc Leu Tyr Phe Pro Pro Gly Val Tyr Trp Met Asn Glu Asp Thr Ser Gly Asn Pro Gly
	ageteggeteaaateatatgeggetggateeeaatacetactgggteeatetageeeeag Lys Leu Gly Ser Asn His Met Arg Leu Asp Pro Asn Thr Tyr Typ Val His Leu Ala Pro
	gagcctatgtgaaaggagccattgagtatttcacgaagcaaaatttctatgcaacgggtc Gly Ala Tyr Val Lys Gly Ala Ile Glu Tyr Phe Thr Lys Gln Asn Phe Tyr Ala Thr Gly
	atggcgttctctcaggtgagaactatgtttatcaggccaatgcagctgataactactatg His Gly Val Leu Ser Gly Glu Asn Tyr Val Tyr Gin Ala Asn Ala Ala Asn Asn Tyr Ty
	ccgtcaagagtgatggcacaagcttgagaatgtggtggcacaacaaccttggaggcggtc Ala Val Lys Ser Asp Gly Thr Ser Leu Arg Met Trp Trp His Asn Asn Leu Gly Gly Gly
	aaacatggttttgcatggggcccaccattaatgcaccgccgtttaatacgatggacttca Gln Thr Trp Phe Cys Met Gly Pro Thr Ile Asn Ala Pro Pro Phe Asn Thr Met Asp Phe
	acggaaactctaatatttccagccggattagtgactataagcaggttggcgcttattttt Asn Gly Asn Ser Asn Ile Ser Ser Arg Ile Ser Asp Tyr Lys Gln Val Gly Ala Tyr Phe
	tccaaacagacggaccggagatctacgaggacagtgttgtccatgacgtcttctggcatg Phe Gln Thr Asp Gly Pro Glu Ile Tyr Glu Asp Ser Val Val His Asp Val Phe Trp His
1321	ttaatgatgatgccatcaagacatattattccggagcttcaatttcacgagcaaccatct

- 427 Val Asn Asp Asp Ala Ile Lys Thr Tyr Tyr Ser Gly Ala Ser Ile Ser Arg Ala Thr Ile
- 1381 ggaagtgtcacaatgacccgatcatacagatgggctggacgtcacgaaatctcaccggaa
 447 Trp Lys Cys His Asn Asp Pro Ile Ile Gln Met Gly Trp Thr Ser Arg Asn Leu Thr Gly
- 1441 tcagcattgataacctgcacgtcatccacacgagatatttcaaatctgaaacagtggttc
 467 Ile Ser Ile Asp Asn Leu His Val Ile His Thr Arg Tyr Phe Lys Ser Glu Thr Val Val
- 1501 cttcagcaatcattggagcgtctccattctacgcaagtggaatgactgttgatcccagcg
 487 Pro Ser Ala Ile Ile Gly Ala Ser Pro Phe Tyr Ala Ser Gly Met Thr Val Asp Pro Ser
- 1561 agtccatcagcatgaccatctctaacgtggtgtgtgagggtctatgccctcactgttcc 507 Glu Scr Ilc Scr Met Thr Ile Ser Asn Val Val Cys Glu Gly Leu Cys Pro Ser Leu Phe
- 1621 gtatcactccgcttcagagctacaacaaccttgttgtcaagaacgtggcctttcccgatg
 527 Arg Ile Thr Pro Leu Gin Ser Tyr Asn Asn Leu Val Val Lys Asn Val Ala Phe Pro Asp
- 1681 gactgcagacaaatccaatcggaataggagagagcattataccagcagcttccggctgta 547 Gly Leu Gln Thr Asn Pro Ile Gly Ile Gly Glu Ser Ile Ile Pro Ala Ala Ser Gly Cys
- 1741 caatggacttggaaatcacaaactggaccgtcaaaggacaaaaagtcaccatgcaaaact
 567 Thr Met Asp Leu Glu Ile Thr Asn Trp Thr Val Lys Gly Gln Lys Val Thr Met Gln Asn
- 1801 ttcagtccgggtcacttggccagttcgatatcgatggttcatactggggtcaatggtcca
 587 Phe Gln Ser Gly Ser Leu Gly Gln Phe Asp Ile Asp Gly Ser Tyr Trp Gly Gln Trp Ser
- 1861 taaac<u>taa</u>agctattcccattcacctgagtattttcgtgggttcaatgagttcttgttac
- 1921 tgatggggcccttgctagtggtaaaagtagagggacttgtcctcgccgggcgccaaggaa
- 2041 aaaaaaaaaaa 2052

FIG. 1b

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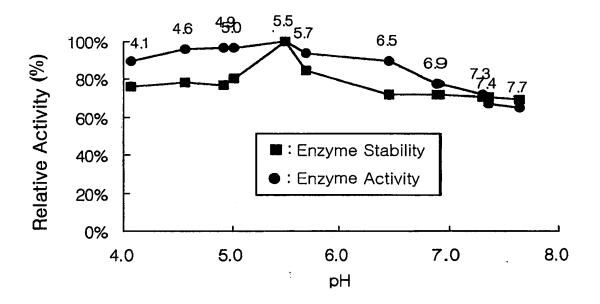


FIG. 2

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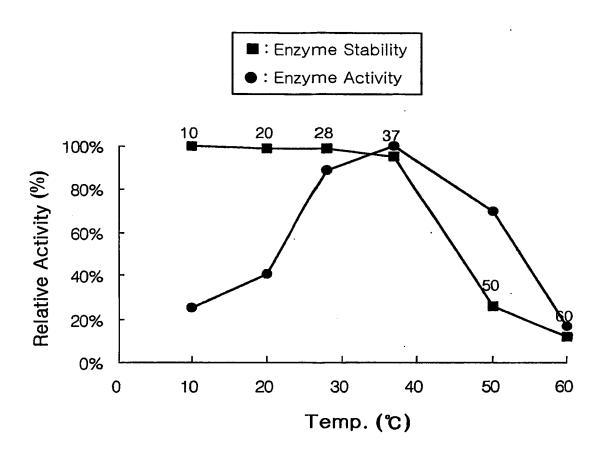


FIG. 3

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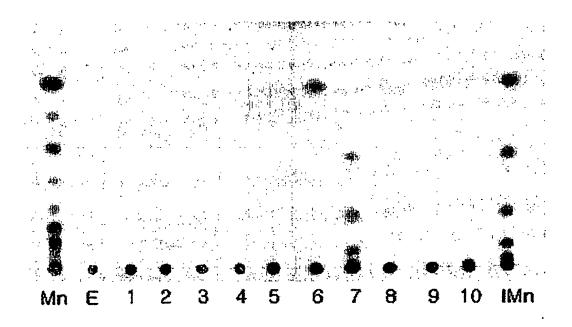


FIG. 4

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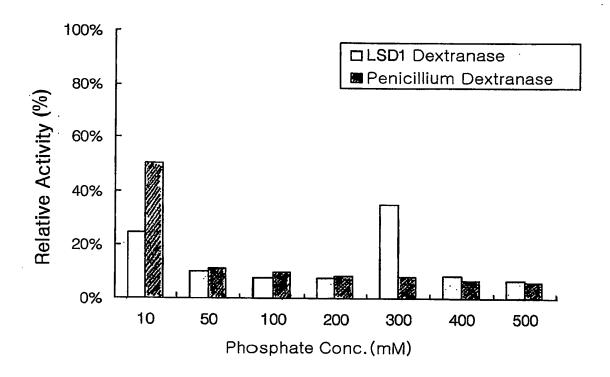


FIG. 5